

IN THE SPECIFICATION

Please replace the paragraph on page 8, lines 1-14 with the following replacement paragraph:

In all these cases, the (replicating or pseudoreplicating) recombinant adenoviruses according to the invention retain the left ITR sequences which are essential for replication and for activating transcription (4 repeated GGTCA motifs located between positions 62 and 99 in the Cav2 genome) as well as the 5'TTGN₈CG-3' type A_I (positions 207-219) and 5'-TTTA/G-3' type A_{II} to A_{IX} encapsidation signals (respectively located at positions 197-200 (A_{II}), 206-209 (A_{III}), 213-216, ~~226-232~~, 228-232 (A_{IV}), 239-242 (A_V), 250-253 (A_{VI}), 258-261 (A_{VII}), 272-275 (A_{VIII}) and 306-309 (A_{IX}) in the Cav2 genome). They also retain all of the E1A coding sequence as well as regions of the E1 gene located downstream thereof (E1A polyadenylation signal and E1B region).

Please replace the Sequence Listing with the attached Substitute Sequence Listing.